

FRET-based nucleosomes sliding

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An abbreviated version of this protocol was published in Nature Communications in Nov 2020

Functional characterization of SMARCA4 variants identified by targeted exome-sequencing of 131,668 cancer patients

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Related files



Epidyne Fret Assay-2.docx



How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Yauch, R. and Fernando, T. (2021). FRET-based nucleosomes sliding. Bio-protocol Preprint. bio-protocol.org/prep1203.
2. Fernando, T. M., Piskol, R., Bainer, R., Sokol, E. S., Trabucco, S. E., Zhang, Q., Trinh, H., Maund, S., Kschonsak, M., Chaudhuri, S., Modrusan, Z., Januario, T. and Yauch, R. L. (2020). Functional characterization of SMARCA4 variants identified by targeted exome-sequencing of 131,668 cancer patients. Nature Communications 11. DOI: [10.1038/s41467-020-19402-8](https://doi.org/10.1038/s41467-020-19402-8)

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